



PCT

## RAW SEQUENCE LISTING

DATE: 07/16/2004

PATENT APPLICATION: US/10/501,040

TIME: 10:15:58

Input Set : A:\82275-sequence.txt

Output Set: N:\CRF4\07162004\J501040.raw

3 <110> APPLICANT: Epigenomics AG  
 5 <120> TITLE OF INVENTION: Method for detecting cytosine-methylation patterns by  
 exponential  
 6 ligation of hybridised probe oligo-nucleotides (MLA)  
 8 <130> FILE REFERENCE: PCT/DE03/00073  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/501,040  
 C--> 10 <141> CURRENT FILING DATE: 2004-07-08  
 10 <160> NUMBER OF SEQ ID NOS: 15  
 12 <170> SOFTWARE: PatentIn version 3.1  
 14 <210> SEQ ID NO: 1  
 15 <211> LENGTH: 33  
 16 <212> TYPE: DNA  
 17 <213> ORGANISM: artificial sequence  
 19 <220> FEATURE:  
 20 <223> OTHER INFORMATION: bisulfite-treated  
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 63 <211> LENGTH: 14  
 64 <212> TYPE: DNA



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**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/501,040

DATE: 07/16/2004

TIME: 10:15:59

Input Set : A:\82275-sequence.txt

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L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date